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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=8; day=27; hr=16; min=56; sec=30; ms=742;]

=====

Reviewer Comments:

<400> 27

Trp Leu Asp Phe

1

1

Please remove the "1" above, which appears at the end of the submitted
file.

Application No: 10587529

Version No: 1.0

Input Set:

Output Set:

Started: 2008-08-23 06:05:24.521

Finished: 2008-08-23 06:05:27.929

Elapsed: 0 hr(s) 0 min(s) 3 sec(s) 408 ms

Total Warnings: 25

Total Errors: 6

No. of SeqIDs Defined: 27

Actual SeqID Count: 27

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
E 257	Invalid sequence data feature in <221> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
E 257	Invalid sequence data feature in <221> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
W 213	Artificial or Unknown found in <213> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)
W 213	Artificial or Unknown found in <213> in SEQ ID (12)
W 213	Artificial or Unknown found in <213> in SEQ ID (13)
W 213	Artificial or Unknown found in <213> in SEQ ID (14)
W 213	Artificial or Unknown found in <213> in SEQ ID (15)
W 213	Artificial or Unknown found in <213> in SEQ ID (16)
W 213	Artificial or Unknown found in <213> in SEQ ID (17)
W 213	Artificial or Unknown found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
W 213	Artificial or Unknown found in <213> in SEQ ID (20)

Input Set:

Output Set:

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Total Warnings: 25
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Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (21)
W 213	Artificial or Unknown found in <213> in SEQ ID (22) This error has occurred more than 20 times, will not be displayed
E 257	Invalid sequence data feature in <221> in SEQ ID (26)
E 257	Invalid sequence data feature in <221> in SEQ ID (27)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (27)

SEQUENCE LISTING

<110> WARATAH PHARMACEUTICALS, INC.
CRUZ, Antonio

<120> METHODS AND COMPOSITIONS USING CD3 AGONISTS

<130> 24492-020 CIP NATL

<140> 10587529

<141> 2008-08-23

<150> 60/584,635

<151> 2004-07-01

<160> 27

<170> PatentIn version 3.3

<210> 1

<211> 37

<212> PRT

<213> Artificial

<220>

<223> Synthesized peptide

<400> 1

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Ser	Ser	Tyr	Leu	Glu	Gly	Gln	Ala	Ala	Lys	Glu	Phe	Ile	Ala	Trp	Leu
			20					25					30		

Val	Lys	Gly	Arg	Gly
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<400> 2

His	Asp	Glu	Phe	Glu	Arg	His	Ala	Glu	Gly	Thr	Phe	Thr	Ser	Asp	Val
1				5					10					15	

Ser	Ser	Tyr	Leu	Glu	Gly	Gln	Ala	Ala	Lys	Glu	Phe	Ile	Ala	Trp	Leu
			20					25					30		

Val Lys Gly Arg
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<222> (36)..(36)
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His Asp Glu Phe Glu Arg His Ala Glu Gly Thr Phe Thr Ser Asp Val
1 5 10 15

Ser Ser Tyr Leu Glu Gly Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu
20 25 30

Val Lys Gly Arg
35

<210> 4
<211> 31
<212> PRT
<213> Artificial

<220>
<223> Synthesized Peptide

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His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
1 5 10 15

Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg Gly
20 25 30

<210> 5
<211> 30
<212> PRT
<213> Artificial

<220>

<223> Synthesized Peptide

<400> 5

His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
1 5 10 15

Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg
20 25 30

<210> 6

<211> 30

<212> PRT

<213> Artificial

<220>

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<220>

<221> MOD_RES

<222> (30)..(30)

<223> AMIDATION

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1 5 10 15

Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg
20 25 30

<210> 7

<211> 39

<212> PRT

<213> Heloderma horridum

<300>

<308> P20394

<309> 2004-06-15

<313> (1)..(39)

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His Ser Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Ser
35

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<213> Heloderma suspectum

<300>
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<309> 1997-11-21
<313> (1)..(39)

<400> 8

His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Pro Ser
35

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His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
1 5 10 15

Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Tyr
20 25 30

<210> 10
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<212> PRT
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<220>
<223> Synthesized Peptide

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Asp Leu Ser Lys Gln Met Glu Glu Glu Ala Val Arg Leu Phe Ile Glu

1 5 10 15

Trp Leu Lys Asn Gly Gly Pro Ser Ser Gly Ala Pro Pro Pro Ser
20 25 30

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<222> (1)..(1)
<223> Xaa is pyroglutamate

<400> 11

Xaa Leu Gly Pro Gln Gly Pro Pro His Leu Val Ala Asp Pro Ser Lys
1 5 10 15

Lys Gln Gly Pro Trp Leu Glu Glu Glu Glu Glu Ala Tyr Gly Trp Met
20 25 30

Asp Phe

<210> 12
<211> 34
<212> PRT
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<220>
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<223> Xaa is pyroglutamate

<400> 12

Xaa Leu Gly Pro Gln Gly Pro Pro His Leu Val Ala Asp Pro Ser Lys
1 5 10 15

Lys Gln Gly Pro Trp Leu Glu Glu Glu Glu Glu Ala Tyr Gly Trp Leu
20 25 30

Asp Phe

<210> 13
<211> 17
<212> PRT
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<220>
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<400> 13

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Phe

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Phe

<210> 15
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<223> Synthesized Peptide

<400> 15

Met Gln Arg Leu Cys Val Tyr Val Leu Ile Phe Ala Leu Ala Leu Ala
1 5 10 15

Ala Phe Ser Glu Ala Ser Trp Lys Pro Arg Ser Gln Gln Pro Asp Ala
20 25 30

Pro Leu Gly Thr Gly Ala Asn Arg Asp Leu Glu Leu Pro Trp Leu Glu
35 40 45

Gln Gln Gly Pro Ala Ser His His Arg Arg Gln Leu Gly Pro Gln Gly
50 55 60

Pro Pro His Leu Val Ala Asp Pro Ser Lys Lys Gln Gly Pro Trp Leu
65 70 75 80

Glu Glu Glu Glu Glu Ala Tyr Gly Trp Met Asp Phe Gly Arg Arg Ser
85 90 95

Ala Glu Asp Glu Asn
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<210> 16

<211> 52

<212> PRT

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<220>

<223> Synthesized Peptide

<400> 16

Asp Leu Glu Leu Pro Trp Leu Glu Gln Gln Gly Pro Ala Ser His His
1 5 10 15

Arg Arg Gln Leu Gly Pro Gln Gly Pro Pro His Leu Val Ala Asp Pro
20 25 30

Ser Lys Lys Gln Gly Pro Trp Leu Glu Glu Glu Glu Glu Ala Tyr Gly
35 40 45

Trp Met Asp Phe
50

<210> 17
<211> 14
<212> PRT
<213> Artificial

<220>
<223> Synthesized Peptide

<400> 17

Trp Leu Glu Glu Glu Glu Glu Ala Tyr Gly Trp Met Asp Phe
1 5 10

<210> 18
<211> 6
<212> PRT
<213> Artificial

<220>
<223> Synthesized Peptide

<400> 18

Tyr Gly Trp Met Asp Phe
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<210> 19
<211> 6
<212> PRT
<213> Artificial

<220>
<223> Synthesized Peptide

<400> 19

Tyr Gly Trp Leu Asp Phe
1 5

<210> 20
<211> 31
<212> PRT
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<220>
<223> Synthesized Peptide

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<223> Xaa is Pro or Tyr

<400> 20

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1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Xaa
20 25 30

<210> 21

<211> 40

<212> PRT

<213> Artificial

<220>

<223> Synthesized Peptide

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<221> MISC_FEATURE

<222> (2)..(3)

<223> Xaa is Ser or Asp Xaa is Gly or Phe

<400> 21

His Xaa Xaa Gly Thr Phe Ile Thr Ser Asp Leu Ser Lys Gln Met Glu
1 5 10 15

Glu Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro
20 25 30

Ser Ser Gly Ala Pro Pro Pro Ser
35 40

<210> 22

<211> 44

<212> PRT

<213> Artificial

<220>

<223> Synthesized Peptide

<400> 22

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1 5 10 15

Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
20 25 30

Ser Gly Ala Pro Pro Ser Lys Lys Lys Lys Lys Lys
35 40

<210> 23
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<212> PRT
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<220>
<223> Synthesized Peptide

<400> 23

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<210> 24
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Tyr Gly Trp Leu Asp Phe
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<210> 25
<211> 10
<212> PRT
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<220>
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<400> 25

Gly Ala Gly Ala Gly Ala Gly Ala Gly Ala
1 5 10

<210> 26
<211> 4
<212> PRT
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<220>
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<223> AMIDATION

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Trp Met Asp Phe
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<210> 27
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<223> AMIDATION

<400> 27

Trp Leu Asp Phe
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